

Table S1

Taxon designation of fungal endophytes from roots and leaves of *S. purpurea* based on sequence data from the internal transcribed spacer regions of nuclear ribosomal RNA (ITS rRNA).

Genus (stated in GenBank)	Phylum; Subclass; Order;	Isolates	ITS (No.)*	Most closely related species**	% similarity
<i>Darksidea</i>	Ascomycota; Dothideomycetes; Pleosporales	ZMr06	MK102695	<i>Darksidea delta</i> (KP183980)	97.23
<i>Saccharicola</i>	Ascomycota; Dothideomycetes; Pleosporales	ZMr02	MK102684	<i>Saccharicola bicolor</i> (KT367526)	100.00
<i>Saccharicola</i>	Ascomycota; Dothideomycetes; Pleosporales	ZMr12	MK102685	<i>Saccharicola bicolor</i> (AF455415)	100.00
<i>Saccharicola</i>	Ascomycota; Dothideomycetes; Pleosporales	ZMr21	MK102687	<i>Saccharicola bicolor</i> (KP117300)	100.00
<i>Saccharicola</i>	Ascomycota; Dothideomycetes; Pleosporales	ZMr24	MK102686	<i>Saccharicola bicolor</i> (KP276515)	100.00
<i>Saccharicola</i>	Ascomycota; Dothideomycetes; Pleosporales	ZMr29	MK102688	<i>Saccharicola bicolor</i> (MH087100)	100.00
<i>Saccharicola</i>	Ascomycota; Dothideomycetes; Pleosporales	ZMr35	MK102689	<i>Saccharicola bicolor</i> (KM979977)	100.00
<i>Saccharicola</i>	Ascomycota; Dothideomycetes; Pleosporales	ZMr38	MK102690	<i>Saccharicola bicolor</i> (JX981466)	99.35
<i>Saccharicola</i>	Ascomycota; Dothideomycetes; Pleosporales	ZMr48	MK102691	<i>Saccharicola bicolor</i> (KJ188730)	99.58
<i>Stagonospora</i>	Ascomycota; Dothideomycetes; Pleosporales	ZMr30	MK102693	<i>Stagonospora perfecta</i> (KF251258)	98.48
<i>Stagonospora</i>	Ascomycota; Dothideomycetes; Pleosporales	ZMr44	MK102694	<i>Stagonospora perfecta</i> (KF251258)	97.23
<i>Paraphoma</i>	Ascomycota; Dothideomycetes; Pleosporales	ZMr04	MK102697	<i>Paraphoma</i> sp. (KT269100)	99.56
<i>Paraphoma</i>	Ascomycota; Dothideomycetes; Pleosporales	ZMr11	MK102698	<i>Paraphoma</i> sp. (KT269033)	100.00
Undefined genus	Ascomycota; Dothideomycetes; Pleosporales	ZMr23	MK102696	<i>Pleosporales</i> sp. (KT269765)	99.56
<i>Phialophora</i>	Ascomycota; Eurotiomycetes; Chaetothyriales	ZMr28	MK102700	<i>Phialophora mustea</i> (JX145398)	99.81
<i>Cadophora</i>	Ascomycota; Leotiomycetes; Helotiales	ZMr07	MK102699	<i>Cadophora</i> sp. (KT264475)	99.23
<i>Cadophora</i>	Ascomycota; Leotiomycetes; Helotiales	ZMr40	MK102701	<i>Cadophora</i> sp. (KT268419)	98.89
<i>Scytalidium</i>	Ascomycota; Leotiomycetes; Helotiales	ZMr13	MK102670	<i>Scytalidium</i> sp. YG-2010a (HQ213805)	99.35
Undefined genus	Ascomycota; Leotiomycetes; Helotiales	ZMr33	MK102702	<i>Helotiales</i> sp. r339 (HQ649857)	99.84

<i>Trichoderma</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr18	MK102662	<i>Trichoderma citrinum</i> (DQ865099)	92.45
<i>Trichoderma</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr39	MK102663	<i>Trichoderma citrinum</i> (KF887057)	99.39
<i>Trichoderma</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr19	MK102667	<i>Trichoderma</i> sp. (KP714590)	95.89
<i>Verticillium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr25	MK102671	<i>Verticillium leptobactrum</i> (KJ188658)	98.56
<i>Verticillium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr26	MK102672	<i>Verticillium</i> sp. TF17TTW (FJ948142)	97.49
<i>Fusarium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr01	MK102655	<i>Fusarium tricinctum</i> (JQ846085)	97.56
<i>Fusarium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr27	MK102656	<i>Fusarium tricinctum</i> (KF010839)	99.21
<i>Fusarium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr09	MK102659	<i>Fusarium</i> sp. (JQ388254)	100.00
<i>Fusarium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr41	MK102657	<i>Fusarium</i> sp. (JQ388252)	99.13
<i>Fusarium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr42	MK102660	<i>Fusarium</i> sp. (AY924269)	99.78
<i>Fusarium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr47	MK102661	<i>Fusarium</i> sp. (HQ130669)	99.45
<i>Fusarium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr05	MK102658	<i>Fusarium</i> sp. (KT270287)	99.49
<i>Myrothecium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr37	MK102665	<i>Myrothecium inundatum</i> (JQ936267)	99.59
<i>Stachybotrys</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr22	MK102664	<i>Stachybotrys bisbyi</i> (KP256004)	99.53
<i>Stachybotrys</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMr14	MK102666	<i>Stachybotrys</i> sp. HZ-23 (EU301653)	99.54
Pyricularia	Ascomycota; Sordariomycetes; Magnaporthales	ZMr10	MK102669	<i>Pyricularia ctenantheicola</i> (KM484878)	99.79
<i>Slopeiomyces</i>	Ascomycota; Sordariomycetes; Magnaporthales	ZMr08	MK102668	<i>Slopeiomyces cylindrosporu</i> (JF508361)	98.46
<i>Monosporascus</i>	Ascomycota; Sordariomycetes; Sordariales	ZMr17	MK102683	<i>Monosporascus ibericus</i> (JQ973832)	99.47
<i>Microdochium</i>	Ascomycota; Sordariomycetes; Xylariales	ZMr03	MK102681	<i>Microdochium bolleyi</i> (KT692593)	98.12
<i>Microdochium</i>	Ascomycota; Sordariomycetes; Xylariales	ZMr15	MK102679	<i>Microdochium bolleyi</i> (KC989068)	96.41
<i>Microdochium</i>	Ascomycota; Sordariomycetes; Xylariales	ZMr16	MK102676	<i>Microdochium bolleyi</i> (JQ658340)	99.45
<i>Microdochium</i>	Ascomycota; Sordariomycetes; Xylariales	ZMr31	MK102678	<i>Microdochium bolleyi</i> (AM502264)	99.78
<i>Microdochium</i>	Ascomycota; Sordariomycetes; Xylariales	ZMr32	MK102682	<i>Microdochium bolleyi</i> (KT692593)	99.56
<i>Microdochium</i>	Ascomycota; Sordariomycetes; Xylariales	ZMr34	MK102677	<i>Microdochium bolleyi</i> (MH858255)	99.37
<i>Microdochium</i>	Ascomycota; Sordariomycetes; Xylariales	ZMr36	MK102673	<i>Microdochium bolleyi</i> (KP859010)	99.69
<i>Microdochium</i>	Ascomycota; Sordariomycetes; Xylariales	ZMr43	MK102674	<i>Microdochium bolleyi</i> (KJ188678)	99.41

<i>Microdochium</i>	Ascomycota; Sordariomycetes; Xylariales	ZMr45	MK102680	<i>Microdochium bolleyi</i> (HQ703412)	100.00
<i>Microdochium</i>	Ascomycota; Sordariomycetes; Xylariales	ZMr46	MK102675	<i>Microdochium bolleyi</i> (KF646098)	99.32
<i>Alternaria</i>	Ascomycota; Dothideomycetes; Pleosporales	ZMI09	MK102650	<i>Alternaria carthami</i> (KJ704213)	99.47
<i>Gloeotinia</i>	Ascomycota; Leotiomycetes; Helotiales	ZMI01	MK102637	<i>Gloeotinia</i> sp. LJC-2011(JF461264)	97.48
<i>Heydenia</i>	Ascomycota; Pezizomycetes ;Pezizales	ZMI04	MK102651	<i>Heydenia alpina</i> (JX171178)	95.25
<i>Heydenia</i>	Ascomycota; Pezizomycetes; Pezizales	ZMI13	MK102653	<i>Heydenia myrsines</i> (KU325128)	97.14
<i>Heydenia</i>	Ascomycota; Pezizomycetes; Pezizales	ZMI14	MK102654	<i>Heydenia alpina</i> (KU325118)	97.86
<i>Heydenia</i>	Ascomycota; Pezizomycetes; Pezizales	ZMI15	MK102652	<i>Heydenia arietina</i> (KF574887)	99.25
<i>Simplicillium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMI06	MK102638	<i>Simplicillium chinense</i> (KP034998)	97.47
<i>Fusarium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMI03	MK102645	<i>Fusarium</i> sp. MPG-5 (KC894839)	99.56
<i>Fusarium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMI07	MK102641	<i>Fusarium acuminatum</i> (KT070871)	99.47
<i>Fusarium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMI17	MK102640	<i>Fusarium pseudograminearum</i> (KP726905)	100.00
<i>Fusarium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMI10	MK102642	<i>Fusarium tricinctum</i> (JX406512)	99.47
<i>Fusarium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMI11	MK102643	<i>Fusarium tricinctum</i> (KU350729)	99.18
<i>Fusarium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMI12	MK102644	<i>Fusarium tricinctum</i> (KJ598865)	99.19
<i>Fusarium</i>	Ascomycota; Sordariomycetes; Hypocreales	ZMI16	MK102639	<i>Fusarium tricinctum</i> (KJ598867)	95.15
<i>Chaetomium</i>	Ascomycota; Sordariomycetes; Sordariales	ZMI08	MK102648	<i>Chaetomium murorum</i> (JQ946413)	98.89
<i>Chaetomium</i>	Ascomycota; Sordariomycetes; Sordariales	ZMI18	MK102649	<i>Chaetomium murorum</i> (KT192199)	97.48
<i>Thielavia</i>	Ascomycota; Sordariomycetes; Sordariales	ZMI05	MK102647	<i>Thielavia hyalocarpa</i> (JQ781752)	97.59
Undefined genus	Ascomycota;	ZMI02	MK102646	<i>Ascomycota</i> sp. (KP698348)	96.19

*ITS rDNA sequences of cultural endophytic fungi were deposited at GenBank;

**Matches of ITS rDNA sequences published were also from GenBank.

Table S2

The distribution of fungal taxa and isolate number at species, order, and class level.

Class type (5) and isolate number	Order type (8) and isolate number	OTUs(33) at species level and isolate number
Dothideomycetes(71)	Pleosporales (71)	<i>Alternaria carthami</i> (2), <i>Darksidea delta</i> (5), <i>Paraphoma</i> sp.(3), <i>Pleosporales</i> sp. (1), <i>Saccharicola bicolor</i> (58), <i>Stagonospora perfecta</i> (2), <i>Phialophora mustea</i> (9),
Eurotiomycetes(9)	Chaetothyriales(9)	<i>Cadophora</i> sp.(3), <i>Gloeotinia</i> sp. (5), <i>Helotiales</i> sp. (1), <i>Scytalidium</i> sp. (3),
Leotiomycetes(12)	Helotiales(12)	<i>Heydenia alpina</i> (2), <i>Heydenia myrsinea</i> (14), <i>Heydenia arietina</i> (18),
Pezizomycetes(34)	Pezizales(34)	<i>Fusarium acuminatum</i> (4), <i>Fusarium pseudograminearum</i> (1), <i>Fusarium</i> sp.(62), <i>Fusarium tricinctum</i> (30), <i>Myrothecium inundatum</i> (1), <i>Simplicillium chinense</i> (2), <i>Stachybotrys bisbyi</i> (3), <i>Trichoderma citrinum</i> (3), <i>Trichoderma</i> sp. (3), <i>Verticillium</i> <i>leptobactrum</i> (10), <i>Verticillium</i> sp.(23) ,
Sordariomycetes(196)	Hypocreales(142)	<i>Pyricularia ctenantheicola</i> (1), <i>Slopeiomyces cylindrosporus</i> (1),
	Magnaporthales(2)	<i>Chaetomium murorum</i> (9), <i>Monosporascus ibericus</i> (1), <i>Thielavia hyalocarpa</i> (5),
	Sordariales(15)	<i>Microdochium bolleyi</i> (37),
	Xylariales(37)	Undefined genus,
Ascomycota sp.(1)		

Table S3

The effects of Kruskal-Wallis one-way ANOVA on ranks and Pearson product moment correlation on the endophytic fungal communities of *S. purpurea*.

Samples*	Kruskal-Wallis one-way		Pearson Product	
	ANOVA on ranks		Moment Correlation	
	H	<i>p</i>	CC**	<i>p</i>
Total vs. R	0.564	0.453	0.967	<0.0001
Total vs. L	0.441	0.507	0.283	0.461
R vs. L	0.002	0.964	0.029	0.941

Values in italics indicate significant differences or significant correlations

* R, root; L, leaf

**CC, correlation coefficient

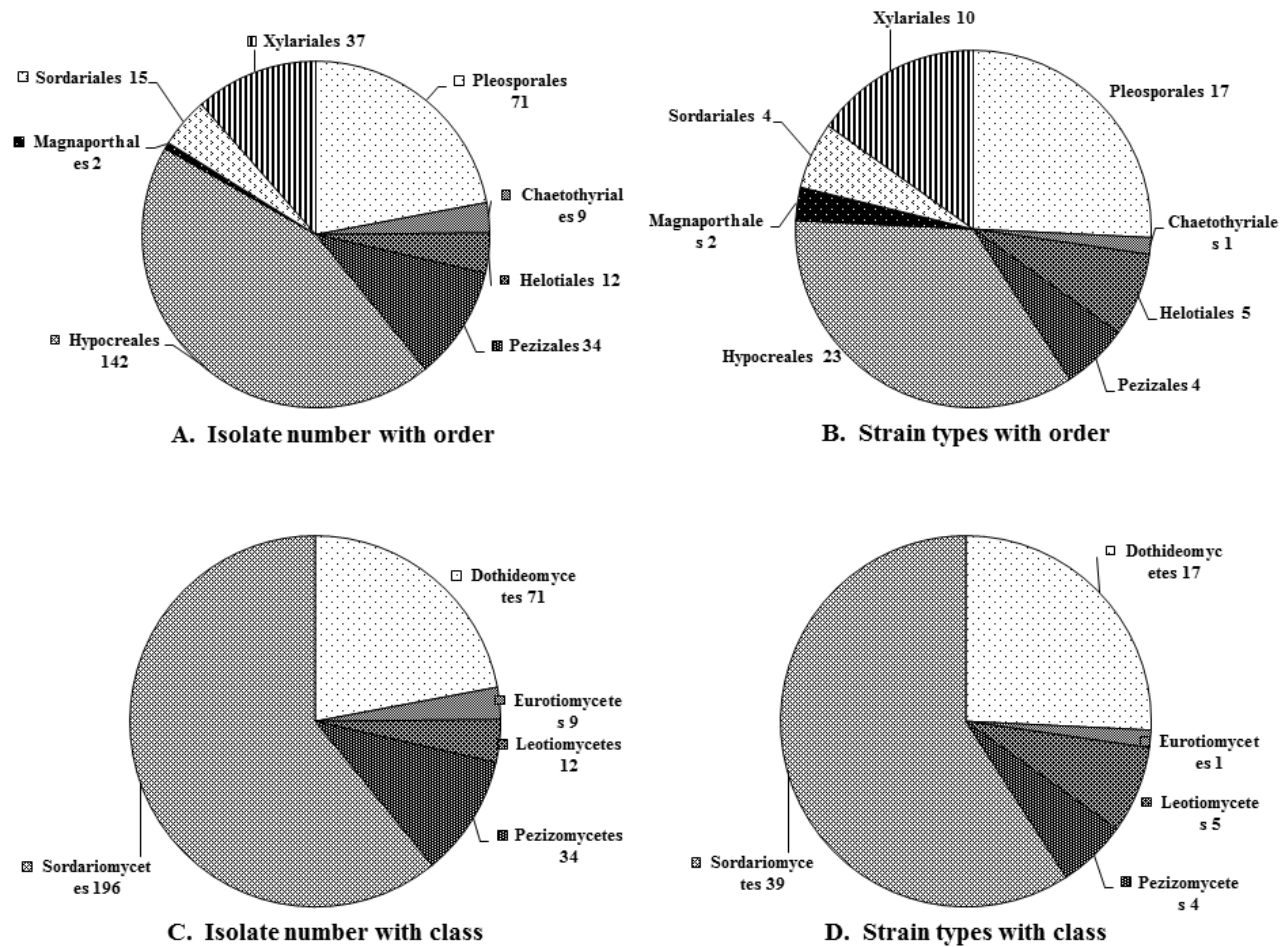


Fig. S1 The distribution of endophytic fungal isolates from *S. purpurea*. The isolate numbers (A) and strain types (B) at the order level; and the isolate numbers (C) and strain types (D) at the class level.